

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/587,389
Source: JFwP
Date Processed by STIC: 8/9/06

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. **EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)**
2. **U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450**
3. **Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314**

Revised 01/10/06

Raw Sequence Listing Error Summary

<u>ERROR DETECTED</u>	<u>SUGGESTED CORRECTION</u>	<u>SERIAL NUMBER:</u> <u>10/587,389</u>
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
1 <input type="checkbox"/> Wrapped Nucleic Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."	
2 <input type="checkbox"/> Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3 <input type="checkbox"/> Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters , instead.	
4 <input checked="" type="checkbox"/> Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.	
5 <input type="checkbox"/> Variable Length	Sequence(s) <input type="checkbox"/> contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6 <input type="checkbox"/> PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) <input type="checkbox"/> . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.	
7 <input type="checkbox"/> Skipped Sequences (OLD RULES)	Sequence(s) <input type="checkbox"/> missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.	
8 <input type="checkbox"/> Skipped Sequences (NEW RULES)	Sequence(s) <input type="checkbox"/> missing. If intentional , please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
9 <input type="checkbox"/> Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.	
10 <input type="checkbox"/> Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence. (see item 11 below)	
11 <input type="checkbox"/> Use of <220>	Sequence(s) <input type="checkbox"/> missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of Sequence Rules	
12 <input type="checkbox"/> PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	
13 <input type="checkbox"/> Misuse of n/Xaa	"n" can only represent a single nucleotide ; "Xaa" can only represent a single amino acid	



IFWP

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/587,389

DATE: 08/09/2006
TIME: 10:06:36

Input Set : N:\SSLM\10587389.txt
Output Set: N:\CRF4\08092006\J587389.raw

4 <110> APPLICANT: Stefano Colloca
5 Alfredo Nicosio
6 Elisabetta Sproreno
7 Agostino Cirillo
8 Bruno Bruni Ercole
9 Annalisa Meola
11 <120> TITLE OF INVENTION: CHIMPANZEE ADENOVIRUS VACCINE CARRIERS
14 <130> FILE REFERENCE: ITR0048YP
C--> 16 <140> CURRENT APPLICATION NUMBER: US/10/587,389
C--> 16 <141> CURRENT FILING DATE: 2006-07-25
16 <150> PRIOR APPLICATION NUMBER: 60/538,799
17 <151> PRIOR FILING DATE: 2004-01-23
19 <150> PRIOR APPLICATION NUMBER: PCT/EP2005/000558
20 <151> PRIOR FILING DATE: 2004-01-18
22 <160> NUMBER OF SEQ ID NOS: 125
24 <170> SOFTWARE: FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

1294 <210> SEQ ID NO: 3
1295 <211> LENGTH: 36606
1296 <212> TYPE: DNA
1297 <213> ORGANISM: Chimpanzee Pan 6 (CV32) Genomic
1299 <400> SEQUENCE: 3
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1301 atttggggag ggaggaagggt gattggctgc gggagcggcg accgttaggg gcggggcg 120
1302 tgacgttttgc atgacgtggc tatgaggcg gacgtttttgc caagttctcg tggaaaagt 180
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1304 ggaaaatgagg tggctctggg cggatgcaag tggaaaacggg ccattttcgc gggaaaactg 300
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1306 ggccgagtag acttgaccg attacgtggg ggtttcgatt accgtatttt tcacctaatt 420
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1316 aggaggcgat tcgagctgcg gtgaaccagg gagtgaaaac tgcggcgag agcttagcc 1020
1317 tggactgtcc tactctgccc ggacacggct gtaagtcttgc tgaatttcat cgcatgaata 1080

Does Not Comply
Corrected Diskette Needed

see item 4 on
End Summary Sheet
sup. 214-6, 9-12, 14, 15,
17-18

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Input Set : N:\SSLM\10587389.txt
Output Set: N:\CRF4\08092006\J587389.raw

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 1321 tcagagtgc a tttcatcacc cccagaaatt ggcaggaac cggccgaaga tattattcat 1320
 1322 agaccagttg cagttagagt caccggccgg agagcagctg tggagagttt ggatgacttg 1380
 1323 ctacagggtg gggatgaacc tttggactt tgcgtccggaa aacgccccag gcactaagtg 1440
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 1325 atccgtgttgc actttaagtg cgtgtttt gactcagggg tggggactgt gggtatataa 1560
 1326 gcagggtcag acctgtgtgg tcagttcaga gcaggactca tggagatctg gactgtctt 1620
 1327 gaagacttc accagactag acagttgcta gagaactcat cggaggaggctt ctcttacctg 1680
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 1330 ggccatcagt ctcactttaa ccagagtatt ctgagagccc ttgactttt tactcctggc 1860
 1331 agaactacccg ccgcggtagc ctttttgcc ttttatttctt acaaattggag tcaagaaacc 1920
 1332 catttcagca gggattaccg tctggactgc ttagcagtag ctttggag aacatggagg 1980
 1333 tgccagcgc tgaatgcaat ctccggctac ttggccagtc agccggtaga cacgctgagg 2040
 1334 atcctgagtc tccagtcacc ccaggaacac caacgcccgc a c a g c c g c a g c a g 2100
 1335 cagcaagagg aggaccgaga agagaacccg agagccggc tggaccctcc ggtggcggag 2160
 1336 gaggaggagt agctgactt tttccggc tgcggccgggt gctgactagg tcttccagtg 2220
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 1339 agggggataga tgaggctctg gtgtgcattt a g a a a t t t t c c t a g a a c a a a g t c a a g a c t t 2400
 1340 gttgggttgc gcccgggat gattggggagg tagccatcag gaattatgcc a a g c t g g c t c 2460
 1341 tgaagccaga caagaagtac a a g a t t a c c a a a c t g a t t a a t a t c a g a a a t t c t c t c t a c a 2520
 1342 tttcaggggaa tggggccgag gtggagatca gtacccagga gagggtggcc ttcagatgtt 2580
 1343 gtatgatgaa tatgtacccg ggggtgggg gcatggagg g a t c a c t t t a t g a a c a c g a 2640
 1344 gtttcagggg tgatgggtat aatgggggg tctttatggc caacaccaag ctgacagtgc 2700
 1345 acggatgctc cttctttggc ttcaataaca tgcgtcatca ggcctgggc a g t g t t c a g 2760
 1346 tgagggatg cagctttca gccaacttgg tgggggtcg gggcagaacc a a g a g c a a g g 2820
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 1349 cccaaagtcaa gcataacatg atctgtgggg cctcggatga g c g c g g c t a c a g a t g c t g a 3000
 1350 cctgcgcggg tggggacagc catatgtgg ccaccgtgca tgcgtccctcg c a c c c c c g c a 3060
 1351 agacatggcc cgaggctcgag cacaacgtca tgacccctgg caatgtc a c t g g c t c c c 3120
 1352 gcccggcat gttcatgc taccagtgc a catgcaatt tgcgtgggtg ctgctggagc 3180
 1353 ccgtatgcgt gtccagatgt agcctgacgg ggggttttgc a t g a a t g t g g a g c t g t g g a 3240
 1354 aaattctgag atatgatgaa tccaaagacca ggtgcggggc tgcgtatgc ggaggcaga 3300
 1355 acggccaggct tcagccctgt tgcgtggagg tgacggaggc cctgcgaccc gatcatttgg 3360
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 1357 agtgtttggg gctgggtgtt ggcctgcatg a g g g g c a g a a t g a c t a a a a t t c t g g t t t t 3480
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 1360 tggacggccg gcccgtgcag cccgcgaact ctcaaccct gacctacgcg acctgagct 3660
 1361 cctcgccgtt ggacgcagct gcccggccag ctgcgtctt cggccgcagc gccgtgcgcg 3720
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 1365 cgggttgcac ggtgaaaacc aaataaaaaa tgaatcaata a a t a a a c c g a g a c g g t t g t t 3960
 1366 gattttaca cagagtctt aatcttatt tgattttcg cgcgcggtag g c c c t g g a c c 4020

see p. 18
for error
explanation

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/587,389

DATE: 08/09/2006
TIME: 10:06:36

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Output Set: N:\CRF4\08092006\J587389.raw

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DATE: 08/09/2006
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Input Set : N:\SSLM\10587389.txt
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 1664 attcataata aacagcacat gtttatgccca ccttctctga ggctctgact ttatTTAGAA 21840
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Output Set: N:\CRF4\08092006\J587389.raw

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 1713 gacgggcata ggcgtgtggc agcagtgcct ggaggagcag aacctgaaag agctctgca 24780
 1714 gtcctgcag aagaacctca aggcctgtg gaccgggtc gacgagcgta ccaccgcctc 24840
 1715 ggacctggcc gacccatct tccccagcgc ctcgcggctg acgctgcgc acgggctgccc 24900
 1716 cgactttatg agccaaagca tggcaaaaa cttcgcgtt ttcatcctcg aacgctccgg 24960
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 1718 gtgcggcccg cgcgtctggc gccactgcta ctgcgtgcgc ctggccaaact acctggcccta 25080
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 1721 gatcatcgcc acctcgagt tgcacggccc cggcgcacggc gagggcaagg ggggtctgaa 25260
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 1751 tccctcggt tactcaacc ctttctcgg ctcggccgc cactacccgg acgagttcat 27060 *see p.18*
 1752 cccgaacttc gacccatca gcgactcggt ggacggctac gattgaatgt cccatggtg 27120
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 1767 tgggtgc当地 ttgtgggaa aagaaatggg gaagatcacc ctatgtgagct ggggtgc当地 28020
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 1777 aaattttta aacattgtgc tcagttatgc ttatgc当地 ggttatgcaaa acatacagaa 28620
 E--> 1778 aaccctttat gttagatctg atggtaact agagatacc caatcacaag ccaagggtgc 28680 *see p. 18*
 1779 atggtaattt tataaaacca acactgatcc agttaaactt tgtaagggtg aattccg 28740
 1780 tacacataaa actccactt catttatgg cagaataat aatcttacac tt当地ttaat 28800
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 1851 ttggactac ccctgaccca agccctaattt gtcaattact ttcagacaga gatgccaat 33000
 1852 ttactctctg tcttacaaaa tgcggtagtc aaataactagg cactgtggca gtggcggctg 33060
 1853 ttactgttagg atcagcacta aatccaattt atgacacagt caaaagcgcctt atagtttcc 33120
 1854 tttagatttgc ttccgatgggt gtactcatgt ccaaactcatc aatggtaggt gattactgg 33180
 1855 acttttaggaa gggacagacc actcaaagtg tagcctatac aaatgctgtg ggattcatgc 33240
 1856 caaatataagg tgcataatcca aaaacccaaa gtaaaacacc taaaatagc atagtcagtc 33300
 1857 aggtatattt aactggagaa actactatgc caatgacact aaccataact ttcaatggca 33360
 1858 ctgatgaaaa agacacaacc ccagtttgc cctactctat gacttttaca tggcagtgg 33420

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/587,389

DATE: 08/09/2006
TIME: 10:06:36

Input Set : N:\SSLM\10587389.txt
Output Set: N:\CRF4\08092006\J587389.raw

1859 ctggagacta taaggacaaa aatattacct ttgctaccaa ctcattctct tttcctaca 33480
 1860 tcgcccagga ataatcccac ccagcaagcc aaccctttt cccaccacct ttgtctatat 33540
 1861 gaaaaactctg aaacagaaaa ataaaagtca agtgtttat tgaatcaaca gttttacagg 33600
 1862 actcgagcag ttatTTTcc tccaccctcc caggacatgg aatacaccac cctctcccc 33660
 1863 cgcacagcct tgaacatctg aatgccattt gtatggaca tgctttggc ctccacgttc 33720
 1864 cacacagttt cagagcgacg cagtctcgga tcggtcaggg agatgaaacc ctccgggac 33780
 1865 tcccgcatct gcacccatca gctcaacagc tgaggattgt cctcggtgg cgggatcacg 33840
 1866 gttatctgga agaagcagaa gagcggcggt gggaaatcata gtccgcgaac ggatcggcc 33900
 1867 ggtgggtctg catcaggccc cgcagcagtc gctccgcgg cgcgtccgtc aagctgctgc 33960
 1868 tcagggggtt cgggtccagg gactccctca gcatgatgcc cacggccctc agatcagtc 34020
 1869 gtctggtgcg gccccgcgcg cagcgcatgc gaatctcgct caggtcaactg cagtaacgtgc 34080
 1870 aacacaggac caccagggtt ttcaacagtc catagttcaa cacgctccag ccgaaactca 34140
 1871 tcgcgggaag gatgctaccc acgtggccgt cgtaccagat cctcaggtaa atcaagtggc 34200
 1872 gctccctcca gaagacgctg cccatgtaca tgatctcctt gggcatgtgg cggttcacca 34260
 1873 cctcccggtt ccacatcacc ctctggtga acatgcagcc cggatgatc ctgcggAAC 34320
 1874 acaggccag caccggcccg cccgcctatgc agcgaagaga ccccgatcc cgcaatgac 34380
 1875 aatggaggac ccaccgctcg taccctgttca tcatctggc gctgaacaag tctatgttg 34440
 1876 cacagcacag gcatatgctc atgcatctct tcagcactct cagctccctg ggggtcaaaa 34500
 1877 ccataccca gggcacgggg aactcttgcg ggacagcgaa ccccgagaa cagggcaatc 34560
 1878 ctgcacata acttacattt tgcatggaca gggatcgca atcaggcagc accgggtgat 34620
 1879 cctccaccag agaagcgcgg gtctcggtct cctcacagcg tggtaagggg gccggccgat 34680
 1880 acgggtgatg gccccgcgcg gctgatcggt ttctcgaccg tgtcatgatc cagttgttt 34740
 1881 cggacatttt cgtactttgt gtagcagaac ctggccggg cgctgcacac cgatcgccgg 34800
 1882 cggcggcttc ggcgttggc acgctcggtg ttaaagtgtt aaaaacagca ctctctcaga 34860
 1883 ccgtgcagca gatctaggcc ctcaggatgc atgaagatcc catcatgcct gatagctctg 34920
 1884 atcacatcga ccacccgttca atggggccagg cccagccaga ttagtgcattt ttgttggtt 34980
 1885 tcggtgacgg cgggggaggg aagaacagga agaaccatga ttaacttttta atccaaacgg 35040
 1886 tctcgagca cttccaaatg aaggtcacgg agatggcacc tctcgcccc gctgtgttg 35100
 1887 tggaaaataa cagccaggc aaagggtata cgggtctcgat gatgttccac ggtggcttcc 35160
 1888 agcaaaagcct ccacgcgcac atcagaaaaca agacaatagc gaaagccggg ggttctcta 35220
 1889 attccctcaac catcatgttca cactccgtca ccattcccccag ataattttca ttttccagc 35280
 1890 cttgaatgtat tcgaacttagt tcctgaggtt aatccaaagcc agccatgata aaaagctcgc 35340
 1891 gcagagcacc ctccacccggc attcttaagc acaccctcat aattccaaaga tattctgctc 35400
 1892 ctggttcacc tgacgcagat tgacaacggg aatataaaaaa tctctccgc gatccctgag 35460
 1893 ctccctccctc agcaataact gtaagtactt tttcatatcg tctccgaaat ttttagccat 35520
 1894 aggaccccca ggaataagag aaggccaaacg cacattacag ataaaccgaa gtcgggggg 35580
 1895 gtgagcatgtt ccaaataatgtt gattgaaaata agcatgtgg cttagccgg tgatatcttc 35640
 1896 cagataactg gacagaaaaat cgggtaaagca atttttaaga aatcaacaa aaaaaaaatc 35700
 1897 ttccaggtgc acgtttaggg cctcgggaaac aacgatggag taagtgcag ggtgcgttc 35760
 1898 cagcatgggtt agtttagctgt tctgtaaaaa aacaaaaaaat aaaaacattaa accatgctag 35820
 1899 cctggcgaac aggtgggtttt atcggtctct ccagcaccag gcaggccacg ggttctccgg 35880
 1900 cgcgaccctc gtaaaaaattt tcgctatgtat tgaaaaccat cacagagaga cttcccccgt 35940
 1901 ggcggcggtt aatgattcga gaagaagcat acaccccccgg gaacatttggc gtcgttgat 36000
 1902 gaaaaaaaaagc ggccgaggaa gcaatgaggc actacaacgc tcactctcaa gtcagcaaa 36060
 1903 gcgatgccc gcgatgttca gacaaaaattt tcaggtgcgtt aaaaaatgtt attactcccc 36120
 1904 tcctgcacag gcagcgaagc tcccgatccc tccagatata catacaacgc ctcagcgtcc 36180
 1905 atagcttacc gaggccgcgcg agcagcgcac cacaacaggc gcaagagtca gaaaaagac 36240
 1906 tgagctctaa cctgtccgccc cgctctctgc tcaatata gccccagatc tacactgac 36300
 1907 taaaggccaa agtctaaaaa taccggccaa ataatcacac acggccagca cacggccaga 36360

RAW SEQUENCE LISTING
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DATE: 08/09/2006
TIME: 10:06:36

Input Set : N:\SSLM\10587389.txt
Output Set: N:\CRF4\08092006\J587389.raw

1908 aaccgggtac acactcgaaa aaatacgcgc acttcctcaa acggccaaac tgccgtcatt 36420
 1909 tccgggttcc cacgctacgt catcaaaaaca cgacttcaa attccgtcga ccgttaaaaa 36480
 1910 catcaccgc cccgccccta acggtcgccg ctcccgacg caatcacctt cctccctccc 36540
 1911 caaattcaaa cagtcattt gcatattaaac ggcacccaaa agtttgaggt atattattga 36600
 1912 tgatgg 36606
 3852 <210> SEQ ID NO: 24
 3853 <211> LENGTH: 2883
 3854 <212> TYPE: DNA
 3855 <213> ORGANISM: Chimpanzee Adenovirus- ChAd 17 Hexon
 3857 <400> SEQUENCE: 24
 3858 atacgcaccc catcgatgat gccgcagtgg tcgtacatgc acatctcggg ccaggacgcc 60
 E--> 3859 ~~t~~tgagtacc tgagccccgg gctgggtcag ttcgccccgg ccacccgagag ctacttcagc 120 *sep 18*
 3860 ctgagaaca agtttagaa cccacgggtg ggcgcacgc acgatgtgac caccgaccgg 180
 3861 tctcagcgcc tgacgctgct gttcattccc gtggacccggc aggacacccgc gtactcgtac 240
 3862 aaggcgcggt tcaccctggc cgtggggcgac aaccgcgtgc tgacatggc ctccacctac 300
 3863 tttgacatcc gcgggggtct ggaccgggggt cccacttca agccctactc tggcaccggcc 360
 3864 tacaactccc tggcccccgggaa gggcgctccc aactccgtcg agtggggagca agagggaaact 420
 3865 caggcagttg aagaagcagc agaagaggaa gaagaagatg ctgacgggtca agctgaggaa 480
 3866 gagcaagcag ctacaaaaaa gactcatgta tatgctcagg ctcccccatttc tggcgaaaaaa 540
 3867 attagtaaaag atggtctgca aataggaacg gacgctacag ctacagaaca aaaacctatt 600
 3868 tatgcagacc ctacattcca gcccgaaccc caaatcgggg agtcacagtg gaatgaggca 660
 3869 gatgctacag tgcggcgccgg tagagtgcta aagaaatcta ctcccatgaa accatgctat 720
 3870 ggttcctatg caagaccac aatgctaattt ggaggtcagg gtgtactaacc ggcaaatgcc 780
 3871 cagggacagc tagaatctca gggtgaaatg caattctttt caacttctga aaacgcccgt 840
 3872 aacgagacta acaacattca gcccaatttggt gtcgtgtata gtgaggatgt gcacatggag 900
 3873 acccccggata cgcacccatttc ttacaagccc gcaaaaaagcg atgacaatttcc aaaaatcatg 960
 3874 ctgggtcagc agtccatgcc caacagaccc aattacatcg gttcagaga taactttatc 1020
 3875 ggcctcatgt attacaatag cactggcaac atgggagtgc ttgcagggtca ggcctctcag 1080
 3876 ttgaatgcag tggggactt gcaagacaga aacacagaac tgccttacca gctcttgctt 1140
 3877 gattccatgg gtgacagaac cagatactttt tccatgtgga atcaggcagt ggacagttat 1200
 3878 gaccaggatg ttagaattat tgaaaatcat ggaactgaaag acgagctccc caactattgt 1260
 3879 ttccctctgg gtggcatagg ggtaactgac acttaccagg ctgttaaaac caacaatggc 1320
 3880 aataaacgggg gccaggtgac ttggacaaaaa gatgaaactt ttgcagatcg caatgaaata 1380
 3881 ggggtggaa acaatttgc tatggagata aacctcagtg ccaacctgtg gagaacttc 1440
 3882 ctgtactcca acgtggcgct gtacctacca gacaagctt agtacaaccc ctccaatgtg 1500
 3883 gacatctctg acaaccccaa cactacgtat tacatgaaca agcgagtggt gggccgggg 1560
 3884 ctggtgact gtcacatcaa cctggcgccg cgctggcgc tgactacat ggacaacgtc 1620
 3885 aacccttca accaccacccg caatgcgggc ctgcgttacc gtcctatgtc cctggcaac 1680
 3886 gggcgctacg tgcccttcca catccaggtg ccccaagaatg tctttgcattt caagaacctc 1740
 3887 ctcctcctgc cgggctctta caccctacgag tggaaacttca ggaaggatgt caacatggtc 1800
 3888 ctccagagct ctctggtaa cgatctcagg gtggacgggg ccagcatcaa gttcgagagc 1860
 3889 atctgcctct acgcccaccc ttccccatg gcccacaaca cggcctccac gtcgaggcc 1920
 3890 atgctcagga acgacaccaa cgaccgttcc ttcaatgact acctctccgc cgccaaatcg 1980
 3891 ctctacccca taccggccaa cgccaccaac gtcccccattt ccattccctc ggcacactgg 2040
 3892 gcgcccttcc gggcgtggc cttcacccgc ctcaagacca aggagacccc ctccctggc 2100
 3893 tcgggattcg accccacta caccctactcg ggctccattt cctaccgttca cggcaccttc 2160
 3894 tacctcaacc acactttcaa gaagggtctcg gtcacccctcg actctctcggt cagctggccg 2220
 3895 ggcaacgacc gtctgctcac ccccaacccgat gtcgagatca agcgctcggt cgacggggag 2280
 3896 qqctacaacq tggcccaatgtca aacatqacc aqqactqgt tcctqgttca qatqctqgccc 2340

RAW SEQUENCE LISTING
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Input Set : N:\SSLM\10587389.txt
Output Set: N:\CRF4\08092006\J587389.raw

3897 aactacaaca tcggcttacca gggcttctac atcccagaga gctacaagga caggatgtac 2400
 3898 tccttcttca ggaacttcca gcccatttgcg cggcagggtgg tggaccagac caagtacaag 2460
 3899 gactaccagg aggtgggcat catccaccag cacaacaact cgggcttcgt gggctacctc 2520
 3900 gccccccacca tgcgcgaggg acaggcctac cccgcact tccctatcc gctcataggc 2580
 3901 aagaccgcgg tgcacagcat caccaggaaa aagttctctgcgacccac cctctggcgc 2640
 3902 atcccccttcc cagcaactt catgtccatg ggtgcgtct cggacctggg ccagaacttg 2700
 3903 ctctacgcac actccgccc cggcctcgac atgaccttcg aggtcgaccc catggacgag 2760
 3904 cccacccttc tctatgttct gttcgaagtc tttgacgtgg tccgggtcca ccagccgcac 2820
 3905 cgcggcgtca tcgagaccgt gtacctgcgt acggcccttct cggccggcaa cgccaccacc 2880
 3906 taa 2883
 10532 <210> SEQ ID NO: 125
 10533 <211> LENGTH: 933
 10534 <212> TYPE: PRT
 10535 <213> ORGANISM: Chimpanzee Adenovirus- CV68 Hexon *ref p. 17*
 10537 <400> SEQUENCE: 125
 10538 Met Ala Thr Pro Ser Met Leu Pro Gln Trp Ala Tyr Met His Ile Ala
 10539 1 5 10 15
 10540 Gly Gln Asp Ala Ser Glu Tyr Leu Ser Pro Gly Leu Val Gln Phe Ala
 10541 20 25 30
 10542 Arg Ala Thr Asp Thr Tyr Phe Ser Leu Gly Asn Lys Phe Arg Asn Pro
 10543 35 40 45
 10544 Thr Val Ala Pro Thr His Asp Val Thr Thr Asp Arg Ser Gln Arg Leu
 10545 50 55 60
 10546 Thr Leu Arg Phe Val Pro Val Asp Arg Glu Asp Asn Thr Tyr Ser Tyr
 10547 65 70 75 80
 10548 Lys Val Arg Tyr Thr Leu Ala Val Gly Asp Asn Arg Val Leu Asp Met
 10549 85 90 95
 10550 Ala Ser Thr Tyr Phe Asp Ile Arg Gly Val Leu Asp Arg Gly Pro Ser
 10551 100 105 110
 10552 Phe Lys Pro Tyr Ser Gly Thr Ala Tyr Asn Ser Leu Ala Pro Lys Gly
 10553 115 120 125
 10554 Ala Pro Asn Thr Cys Gln Trp Thr Tyr Lys Ala Asp Gly Glu Thr Ala
 10555 130 135 140
 10556 Thr Glu Lys Thr Tyr Thr Gly Asn Ala Pro Val Gln Gly Ile Asn
 10557 145 150 155 160
 10558 Ile Thr Lys Asp Gly Ile Gln Leu Gly Thr Asp Thr Asp Asp Gln Pro
 10559 165 170 175
 10560 Ile Tyr Ala Asp Lys Thr Tyr Gln Pro Glu Pro Gln Val Gly Asp Ala
 10561 180 185 190
 10562 Glu Trp His Asp Ile Thr Gly Thr Asp Glu Lys Tyr Gly Gly Arg Ala
 10563 195 200 205
 10564 Leu Lys Pro Asp Thr Lys Met Lys Pro Cys Tyr Gly Ser Phe Ala Lys
 10565 210 215 220
 10566 Pro Thr Asn Lys Glu Gly Gln Ala Asn Val Lys Thr Gly Thr Gly
 10567 225 230 235 240
 10568 Thr Thr Lys Glu Tyr Asp Ile Asp Met Ala Phe Phe Asp Asn Arg Ser
 10569 245 250 255
 10570 Ala Ala Ala Ala Gly Leu Ala Pro Glu Ile Val Leu Tyr Thr Glu Asn
 10571 260 265 270

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Input Set : N:\SSLM\10587389.txt
Output Set: N:\CRF4\08092006\J587389.raw

10572 Val Asp Leu Glu Thr Pro Asp Thr His Ile Val Tyr Lys Ala Gly Thr
10573 275 280 285
10574 Asp Asp Ser Ser Ser Ile Asn Leu Gly Gln Gln Ala Met Pro Asn
10575 290 295 300
10576 Arg Pro Asn Tyr Ile Gly Phe Arg Asp Asn Phe Ile Gly Leu Met Tyr
10577 305 310 315 320
10578 Tyr Asn Ser Thr Gly Asn Met Gly Val Leu Ala Gly Gln Ala Ser Gln
10579 325 330 335
10580 Leu Asn Ala Val Val Asp Leu Gln Asp Arg Asn Thr Glu Leu Ser Tyr
10581 340 345 350
10582 Gln Leu Leu Leu Asp Ser Leu Gly Asp Arg Thr Arg Tyr Phe Ser Met
10583 355 360 365
10584 Trp Asn Gln Ala Val Asp Ser Tyr Asp Pro Asp Val Arg Ile Ile Glu
10585 370 375 380
10586 Asn His Gly Val Glu Asp Glu Leu Pro Asn Tyr Cys Phe Pro Leu Asp
10587 385 390 395 400
10588 Ala Val Gly Arg Thr Asp Thr Tyr Gln Gly Ile Lys Ala Asn Gly Thr
10589 405 410 415
10590 Asp Gln Thr Thr Trp Thr Lys Asp Asp Ser Val Asn Asp Ala Asn Glu
10591 420 425 430
10592 Ile Gly Lys Gly Asn Pro Phe Ala Met Glu Ile Asn Ile Gln Ala Asn
10593 435 440 445
10594 Leu Trp Arg Asn Phe Leu Tyr Ala Asn Val Ala Leu Tyr Leu Pro Asp
10595 450 455 460
10596 Ser Tyr Lys Tyr Thr Pro Ala Asn Val Thr Leu Pro Thr Asn Thr Asn
10597 465 470 475 480
10598 Thr Tyr Asp Tyr Met Asn Gly Arg Val Val Ala Pro Ser Leu Val Asp
10599 485 490 495
10600 Ser Tyr Ile Asn Ile Gly Ala Arg Trp Ser Leu Asp Pro Met Asp Asn
10601 500 505 510
10602 Val Asn Pro Phe Asn His His Arg Asn Ala Gly Leu Arg Tyr Arg Ser
10603 515 520 525
10604 Met Leu Leu Gly Asn Gly Arg Tyr Val Pro Phe His Ile Gln Val Pro
10605 530 535 540
10606 Gln Lys Phe Phe Ala Ile Lys Ser Leu Leu Leu Pro Gly Ser Tyr
10607 545 550 555 560
10608 Thr Tyr Glu Trp Asn Phe Arg Lys Asp Val Asn Met Ile Leu Gln Ser
10609 565 570 575
10610 Ser Leu Gly Asn Asp Leu Arg Thr Asp Gly Ala Ser Ile Ser Phe Thr
10611 580 585 590
10612 Ser Ile Asn Leu Tyr Ala Thr Phe Pro Met Ala His Asn Thr Ala
10613 595 600 605
10614 Ser Thr Leu Glu Ala Met Leu Arg Asn Asp Thr Asn Asp Gln Ser Phe
10615 610 615 620
10616 Asn Asp Tyr Leu Ser Ala Ala Asn Met Leu Tyr Pro Ile Pro Ala Asn
10617 625 630 635 640
10618 Ala Thr Asn Val Pro Ile Ser Ile Pro Ser Arg Asn Trp Ala Ala Phe
10619 645 650 655
10620 Arg Gly Trp Ser Phe Thr Arg Leu Lys Thr Lys Glu Thr Pro Ser Leu

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/587,389

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Input Set : N:\SSLM\10587389.txt
Output Set: N:\CRF4\08092006\J587389.raw

10621	660	665	670
10622	Gly Ser Gly Phe Asp Pro Tyr Phe Val Tyr Ser Gly Ser Ile Pro Tyr		
10623	675	680	685
10624	Leu Asp Gly Thr Phe Tyr Leu Asn His Thr Phe Lys Lys Val Ser Ile		
10625	690	695	700
10626	Thr Phe Asp Ser Ser Val Ser Trp Pro Gly Asn Asp Arg Leu Leu Thr		
10627	705	710	715
10628	720	725	730
10629	735	740	745
10630	750	755	760
10631	765	770	775
10632	780	785	790
10633	800	805	810
10634	815	820	825
10635	830	835	840
10636	845	850	855
10637	860	865	870
10638	880	885	890
10639	895	900	905
E--> 10640	910	915	920
10641	925		
10642	Met Arg Gln Gly Gln Pro Tyr Pro Ala Xaa Tyr Pro Tyr Pro Leu Ile		
10643	820	825	830
10644	835	840	845
10645	850	855	860
10646	875	880	885
10647	885	890	895
10648	900	905	910
E--> 10652	His Arg Gly Val Ile Glu Ala Val Tyr Xaa Arg Thr Pro Phe Ser Ala		
10653	915	920	925
10654	Gly Asn Ala Thr Thr		
10655	930		
E--> 10658	- 1 -		

see p. 18

VARIABLE LOCATION SUMMARY DATE: 08/09/2006
PATENT APPLICATION: US/10/587,389 TIME: 10:06:38

Input Set : N:\SSLM\10587389.txt
Output Set: N:\CRF4\08092006\J587389.raw

Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.

Use of n's and/or Xaa's have been detected in the sequence list. Use of <220> to <223> is MANDATORY if n's or Xaa's are present.

use of <220> to <223> is mandatory if n's or Xaa's are present. in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#: 3; N Pos. 3592, 7705, 11272, 11275, 15203, 24396, 27010, 28655, 30744, 31045

Seq#:24; N Pos. 63 / / /

Seq#:125; Xaa Pos. 826,922

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/587,389

DATE: 08/09/2006

TIME: 10:06:38

Input Set : N:\SSLM\10587389.txt

Output Set: N:\CRF4\08092006\J587389.raw

L:16 M:270 C: Current Application Number differs, Replaced Current Application No
L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:1359 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:3
M:340 Repeated in SeqNo=3
L:3859 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:24
L:10640 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:125
M:340 Repeated in SeqNo=125
L:10658 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:125